```
MAR 2 8 2001 8
                           SEQUENCE LISTING
  <110 Marasco, Wayne
        Mhashilkar, Abner
  <120>
          INTRABODY-MEDIATED CONTROL OF IMMUNE REACTIONS
  <130>
          700157-47577C
  <140> 09/522,727
  <141>
         2000-03-10
  <150> PCT/US98/19563
  <151>
         1998-09-18
  <150>
         60/059,339
  <151>
         1997-09-18
  <160>
         55
  <170>
         PatentIn version 3.0
  <210> 1
  <211> 15
  <212> PRT
  <213>
         Homo sapiens
  <400> 1
  Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser
  1
                                                         15
  <210>
         2
  <211>
         15
  <212>
         PRT
  <213>
         Homo sapiens
  <400> 2
  Glu Ser Gly Arg Ser/Gly Gly Gly Gly Ser Gly Gly Gly Ser
                                                         15
  <210>
         3
  <211>
         14
  <212>
         PRT
  <213>
         Homo sapiens
  <400>
  Glu Gly Lys Ser Ser Gly Ser Glu Ser Lys Ser Thr
```

```
<210>
<211>
       15
<212>
       PRT
<213>
       Homo sapiens
<400>
       4
Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Ser Thr Gln
                                     10
<210>
       5
<211>
       14
<212> PRT
<213>
       Homo sapiens
<400>
Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser/Lys Val Asp
                                     10
<210>
       6
<211>
       14
<212>
       PRT
<213>
       Homo sapiens
<400>
      6
Gly Ser Thr Ser Gly Ser Gly Lys Ser Ser Glu Gly Lys Gly
                                     10
<210>
       7
<211>
       15
<212>
       PRT
<213>
       Homo sapiens
<400>
Lys Glu Ser Gly Ser Val Ser Ser Glu Gln Leu Ala Gln Phe Arg
                                     10
                                                         15
<210>
       8
<211>
       16
<212>
       PRT
<213>
       Homo sapiens
<400>
Glu Ser Gly Ser Val Ser Ser Glu Glu Leu Ala Phe Arg Ser Leu Asp
                                     10
<210>
<211>
       35
```

```
<212>
      DNA
<213>
      Homo sapiens
<400> 9
                                                                     35
tttgcggccg ctcaggtgca rctgctcgag tcygg
<210>
      10
<211>
      66
<212>
      DNA
<213>
      Homo sapiens
<400> 10
agatccgccg ccaccgctcc caccacctcc ggagccaccg ccacctgagg tgaccgtgac
                                                                     60
                                                                     66
crkggt
<210> 11
<211> 69
<212>
      DNA
<213>
      Homo sapiens
<400> 11
ggtggcggtg gctccggagg tggtgggagc ggtggcggcg gatctgagct cswgmtgacc
                                                                     60
                                                                     69
cagtctcca
<210>
       12
<211>
       47
<212>
       DNA
<213>
      Homo sapiens
<400>
      12
gggtctagac tcgaggatcc /tattaacgc gttggtgcag ccacagt
                                                                     47
<210>
       13
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
       13
Ser Glu Lys Asp Glu Leu
<210>
       14
<211/2
       59
       DNA
<2/13>
       Homo sapiens
```

```
<400> 14
                                                                    59
gggtctagac tcgaggatcc ttattacagc tcgtcctttt cgcttggtgc agccacagt
<210>
      15
<211>
      24
<212>
      DNA
<213>
      Homo sapiens
<400> 15
                                                                    24
tttaccatgg aacatctgtg gttc
<210>
      16
<211>
      30
<212>
      DNA
<213>
      Homo sapiens
<400> 16
                                                                    30
ttagcgcgct gaggtgaccg tgaccrkggt
<210> 17
<211> 4
<212> PRT
<213>
      Homo sapiens
<400>
     17
Lys Asp Glu Leu
1
<210> 18
<211>
<212> PRT
<213> Homo sapiens
<400> 18
Asp Asp Glu Leu
<210>
       19
<211>
       4
<212>
       PR7
<213> Hømo sapiens
<400>
      19
Asp/Glu Glu Leu
```

```
<210>
       20
<211>
       4
<212>
       PRT
<213>
       Homo sapiens
<400>
       20
Gln Glu Asp Leu
1
<210>
       21
<211>
       4
<212>
       PRT
<213>
       Homo sapiens
<400>
       21
Arg Asp Glu Leu
<210>
       22
<211>
       7
<212>
       PRT
<213>
       Homo sapiens
<400>
       22
Pro Lys Lys Arg Lys Val
<210>
       23
<211>
       7
<212>
       PRT
<213>
       Homo sapiens
<400> 23
Pro Gln Lys Lys Ile/Lys Ser
<210>
       24
<211>
       5
<212>
       PRT
<213>
       Homo sapiens
<400>
       24
Gln Pro/Lys Lys Pro
<210/>
       25
```

```
<211>
       12
<212> PRT
<213>
      Homo sapiens
<400>
       25
Arg Lys Lys Arg Arg Gln Arg Arg Arg Ala His Gln
                                     10
<210>
       26
<211>
       16
<212>
      PRT
<213>
      Homo sapiens
<400>
       26
Arg Gln Ala Arg Arg Asn Arg Arg Arg Arg Trp Arg Slu Arg Gln Arg
                                                          15
                                     10
<210>
       27
<211>
       19
<212>
      PRT
<213>
      Homo sapiens
<400>
       27
Met Pro Leu Thr Arg Arg Pro Ala Ala Ser Gln Ala Leu Ala Pro
                                     10
1
                                                          15
Pro Thr Pro
<210>
       28
<211>
       15
<212>
       PRT
<213> Homo sapiens
<400> 28
Met Asp Asp Gln Arg Asp Leu Ile Ser Asn Asn Glu Gln Leu Pro
                                     10
                                                          15
<210>
       29
<211>
       32
<212>
       PRT
<213>
       Homo sapiens
<220>
<221>,
       UNSURE
<222%
       (7)(8)(32)
<22/3>
       UNSURE
```

```
<400> 29
Met Leu Phe Asn Leu Arg Xaa Xaa Leu Asn Asn Ala Ala Phe Arg His
                                     10
Gly His Asn Phe Met Val Arg Asn Phe Arg Cys Gly Gln Pro Leu Xaa
            20
                                 25
<210>
       30
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
       30
Gly Cys Val Cys Ser Ser Asn Pro
<210>
       31
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
      31
Gly Gln Thr Val Thr Thr Pro Leu
<210>
       32
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
       32
Gly Gln Glu Leu Ser Gln His Glu
<210>
       33
<211>
<212>
       PRT
<213>
       Homo/sapiens
<400>
Gly Asn Ser Pro Ser Tyr Asn Pro
<21Ø>
       34
<21/1>
<212>
       PRT
```

```
<213>
       Homo sapiens
<400>
       34
Gly Val Ser Gly Ser Lys Gly Gln
<210>
       35
<211>
<212>
      PRT
<213>
       Homo sapiens
<400>
       35
Gly Gln Thr Ile Thr Thr Pro Leu
<210>
       36
<211>
<212>
      PRT
<213>
       Homo sapiens
<400>
       36
Gly Gln Thr Leu Thr Thr Pro Leu
<210>
       37
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
       37
Gly Gln Ile Phe Ser Arg Ser Ala
1
<210>
       38
<211>
<212>
       PRT
<213>
       Homo sapiens
<400> 38
Gly Gln /le His Gly Leu Ser Pro
<210 > 39
<211/>
<21/2>
       PRT
<213>
       Homo sapiens
```

```
<400> 39
Gly Ala Arg Ala Ser Val Leu Ser
<210>
      40
<211>
<212> PRT
<213> Homo sapiens
<400>
      40
Gly Cys Thr Leu Ser Ala Glu Glu
<210> 41
<211>
<212> PRT
<213>
      Homo sapiens
<400> 41
Gly Gln Asn Leu Ser Thr Ser Asn
1
<210>
      42
<211>
<212>
      PRT
<213>
      Homo sapiens
<400> 42
Gly Ala Ala Leu Thr Ile/Leu Val
<210> 43
<211> 8
<212>
      PRT
<213> Homo sapjens
<400>
       43
Gly Ala Ala Leu Thr Leu Leu Gly
<210>
<211>
<212>/ PRT
       Homo sapiens
<213/>
<400>
       44
```

```
Gly Ala Gln Val Ser Ser Gln Lys
<210> 45
<211>
<212>
      PRT
<213>
      Homo sapiens
<400>
     45
Gly Ala Gln Leu Ser Arg Asn Thr
<210>
       46
<211> 8
<212> PRT
<213>
      Homo sapiens
<400> 46
Gly Asn Ala Ala Ala Lys Lys
<210>
       47
<211>
      8
<212>
      PRT
<213>
      Homo sapiens
<400> 47
Gly Asn Glu Ala Ser Tyr Pro Leu/
<210>
       48
<211>
<212> PRT
<213> Homo sapiens
<400> 48
Gly Ser Ser Lys Ser Lys Pro Lys
<210> 49
<211>
       38
<212> DNA
<213> Homo sapiens
<400> /49
ccctctagac atatgtgaat tccaccatgg cccaggtc
                                                                  38
```

```
<210>
       50
<211>
       25
<212>
       DNA
<213>
       Homo sapiens
<400>
       50
tgmggagacg gtgaccrwgg tccct
                                                                    25
<210> 51
<211> 837
<212> DNA
<213> human
<220>
<221> CDS
      UNSURE
<222> (1)...(837)
      (505)
<223> UNSURE
<400> 51
atg gaa cat ctg tgg ttc ttc ctt ctc ctg gtg gca gct ccc aga tgg
Met Glu His Leu Trp Phe Phe Leu Leu Leu Val Ala Ala Pro Arg Trp
1
gtc ctg tcc cag gtg caa ctg cag dag/tca ggg gct gag ctg gca aga
Val Leu Ser Gln Val Gln Leu Gln Oxf Ser Gly Ala Glu Leu Ala Arg
            20
                                                     30
cct ggg gct tca gtg aag ttg tck tgc aag gct tct ggc tac acc ttt 144
Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe
        35
act agt cac tgg atg cag tgg gtg aga cag agg cct gga cag ggt ctg 192
Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu
    50
                                             60
gaa tgg att ggg act at/t tat cct gga gat ggt gat act agg tac act 240
Glu Trp Ile Gly Thr /le Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr
65
                    70
                                         75
cag aat ttc aag gg/c aag gcc aca ttg act gca gat aag tcc tcc acc 288
Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr
                                                         95
aca gcc tac thá cac ctc agc agc ttg tca tct gaa gac tct gcg gtc 336
Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val
            100
                                105
                                                     110
tat tat tg/t gca aga gat gag att act acg gtt gta ccc cgg ggg ttt 384
Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe
                            120
                                                 125
gct tax tgg ggc caa ggg acc tcg gtc acc gtc tcc tca ggt ggc ggt 432
Ala Try Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly
    130
                        135
                                             140
gg¢ tcg ggc ggt ggc tcg ggt ggc ggc gga tct gag ctc gtg ctc 480
GYy Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Leu Val Leu
145
                    150
                                         155
                                                             160
acc caa acc cca acc tcc ctg gct ncc tct ctg gga gac aga gtc acc 528
```

```
Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly Asp Arg Val Thr
                165
                                     170
                                                         175
atc agt tgc agg gca agt cag gac att agc agt tat tta aac tgg tat 578/
Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser Tyr Leu Asn Trp Tyr
            180
                                 185
                                                     190
cag cag aaa cca gat gga act att aaa ctc ctg atc tac tac aca tga 624
Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Zer
        195
                             200
                                                 205
aga tta tat tca gga gtc cca cca agg ttc agt ggc agt ggg gét gga 672
Arg Leu Tyr Ser Gly Val Pro Pro Arg Phe Ser Gly Ser Gly Ala Gly
    210
                                             220
                        215
aca gat tat tct ctc acc atc agc aac ctg gag caa gaa gat ttt gcc 720
Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu/Asp Phe Ala
225
                    230
                                         235
                                                              240
act tac ttt tgc caa cag ggt aat gtg att ccg tac acg ttc gga ggg 768
Thr Tyr Phe Cys Gln Gln Gly Asn Val Ile Pro Tyr/Thr Phe Gly Gly
                245
                                     250
                                                         255
ggg acc aag ctg gaa atg aaa cgg gct gat gct gca cca act gta agc 816
Gly Thr Lys Leu Glu Met Lys Arg Ala Asp Ala/Ala Pro Thr Val Ser
            260
                                                     270
                                 265
                                                                  837
gaa aag gac gag ctg taa taa
Glu Lys Asp Glu Leu
        275
<210>
       52
<211>
       277
<212>
       PRT
<213>
       Homo sapiens
<220>
<221>
       UNSURE
<222>
       (169)
<223>
       UNSURE
<400> 52
Met Glu His Leu Tro Phe Phe Leu Leu Leu Val Ala Ala Pro Arg Trp
                                     10
                                                         15
Val Leu Ser Glm Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Ala Arg
                                                     30
                                 25
Pro Gly Ala/Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe
Thr Ser/His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu
Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr
                                                              80
                     70
```

Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe Ala Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Glu/Leu Val Leu Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly/Asp Arg Val Thr Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser/Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Ser Arg Leu Tyr Ser Gly Val Pro Pro Arg/(Ph/e Ser Gly Ser Gly Ala Gly Thr Asp Tyr Ser Leu Thr Ile Ser/Asn Leu Glu Gln Glu Asp Phe Ala Thr Tyr Phe Cys Gln Gln Gly/Asn Val Ile Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Met/Lys Arg Ala Asp Ala Ala Pro Thr Val Ser Glu Lys Asp Glu Leu <210> 53 <211> 837 <212> DNA <213> human <220> <221> CDS <222> (1)/...(837) <400> 53

atg gaa cat ctg tgg ttc ttc ctt ctc ctg gtg gca gct ccc aga tgg

Met glu His Leu Trp Phe Phe Leu Leu Leu Val Ala Ala Pro Arg Trp

10 15 gtc ctg tcc cag gtg caa ctg cag cag tct ggg gct gag ctg aca aga Val Leu Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Thr Arg cct ggg gct tca gtg aag ttg tcc tgc aag gct tct ggc tac acc ttt 1/44 Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe 35 act agt cac tgg atg cag tgg gtg aga cag agg cct gga cag ggt øtg 192 Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly/Leu 50 gaa tgg att ggg act att tat cct gga gat ggt gat act agg /tac act 240 Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr 65 70 80 cag aat ttc aag ggc aag gcc aca ttg act gca gat aag/tcc tcc acc 288 Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr 90 95 aca gcc tac tta cac ctc agc agc ttg tca tct gag gac tct gcg gtc 336 Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser G/Lu Asp Ser Ala Val 100 105 110 tat tat tgt gca aga gat gag att act acg gtx gta ccc cgg ggg ttt 384 Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Yal Val Pro Arg Gly Phe 115 120 125 gct tac tgg ggc caa ggg acc ttg gtc acc gtc tcc tca ggt ggc ggt 432 Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly 130 135 140 ggc tcg ggc ggt ggg tcg ggt \ggc/ggc gga tct gag ctc gtg ctc 480 Gly Ser Gly Gly Gly Ser Gly King Gly Gly Ser Glu Leu Val Leu 145 150 155 160 acc cag tot coa too agt otg tot/goa too ott gga gac aca att acc 528 Thr Gln Ser Pro Ser Ser Leu Sér Ala Ser Leu Gly Asp Thr Ile Thr 165 170 175 atc act tgc cat gcc agt cag aac att aat gtt tgg tta agt tgg tac 576 Ile Thr Cys His Ala Ser Gin Asn Ile Asn Val Trp Leu Ser Trp Tyr 180 185 190 cag cag aaa cca gga aat att cct caa cta ttg atc tat aag gct tcc 624 Gln Gln Lys Pro Gly Agn Ile Pro Gln Leu Leu Ile Tyr Lys Ala Ser 195 200 205 aac ttg cac aca ggc/gtc cca tca agg ttt agt ggc cgt gga tct gga 672 Asn Leu His Thr Gly Val Pro Ser Arg Phe Ser Gly Arg Gly Ser Gly 210 215 220 aca ggt ttc aca/tta acc atc agc agc ctg cag cct gaa gac att ggc 720 Thr Gly Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Gly 225 230 235 240 act tac tac/tgt caa cag ggt caa agt tat cct ctg acg ttc ggt gga 768 Thr Tyr Tyr Cys Gln Gln Gly Gln Ser Tyr Pro Leu Thr Phe Gly Gly 245 250 255 260 ggc acc ag ctg gaa atc aaa cgg gct gat gct gca cca act gta agc 816 Gly Thr/Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala Pro Thr Val Ser 265 275 270 gaa aag gac gag ctg taa taa 837 Glu Lys Asp Glu Leu 280

UN

<210> 54

<211> 277

<212> PRT

<213> Homo sapiens

<400> 54

Met Glu His Leu Trp Phe Phe Leu Leu Leu Val Ala Ala Pro Arg Trp
1 10 15

Val Leu Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Thr Arg 20 25 30

Pro Gly Ala Ser Val Leu Leu Ser Cys Leu Ala Ser Gly Tyr Thr Phe 35

Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu 50 60

Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr 70 75 80

Gln Asn Phe Leu Gly Leu Ala Thr Leu Thr Ala Asp Leu Ser Ser Thr
85 90 95

Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val

Tyr Tyr Cys Ala Arg Asp Glu Ile Whr Thr Val Val Pro Arg Gly Phe 115 120 125

Ala Tyr Trp Gly Gln Gly Thr Zeu Val Thr Val Ser Ser Gly Gly Gly 130 140

Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Glu Leu Val Leu 145 150 150 160

Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly Asp Thr Ile Thr 165 170 175

Ile Thr Cys His Ala Ser Gln Asn Ile Asn Val Trp Leu Ser Trp Tyr
180 185 190

Gln Gln Leu Pro Gly Asn Ile Pro Gln Leu Leu Ile Tyr Leu Ala Ser 195 200 205

Asn Leu His Thr Gly Val Pro Ser Arg Phe Ser Gly Arg Gly Ser Gly 210 220

Thr Gly Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Gly 225 230 235 240

Thr Tyr Tyr Cys Gln Gln Gly Gln Ser Tyr Pro Leu Thr Phe Gly Gly 245 250 Gly Thr Leu Leu Glu Ile Leu Arg Ala Asp Ala Pro Thr Val Ser 260 265 270 Glu Leu Asp Glu Leu 275 <210> 55 <211> 4 <212> PRT <213> Homo sapiens <400> 55 Arg Lys Lys Arg

UN